

Fig. 1A

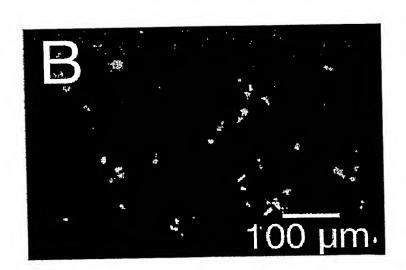


Fig. 1B

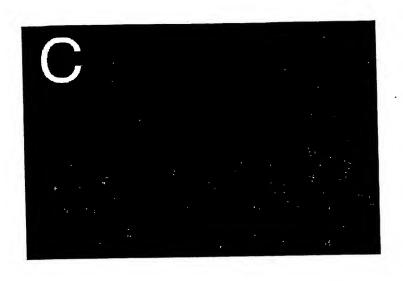


Fig. 10

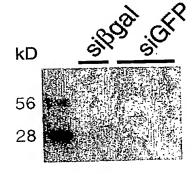


Fig. 1D

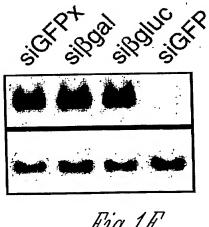


Fig. 1E

siGFP siβgluc

Fig. 1F



Fig. 1G

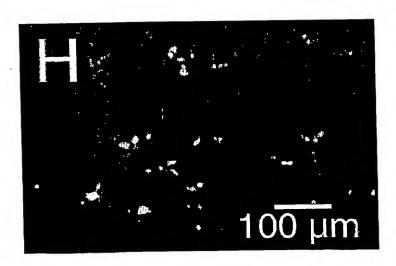


Fig. 1H

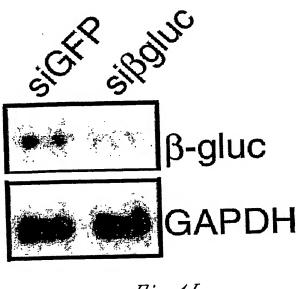
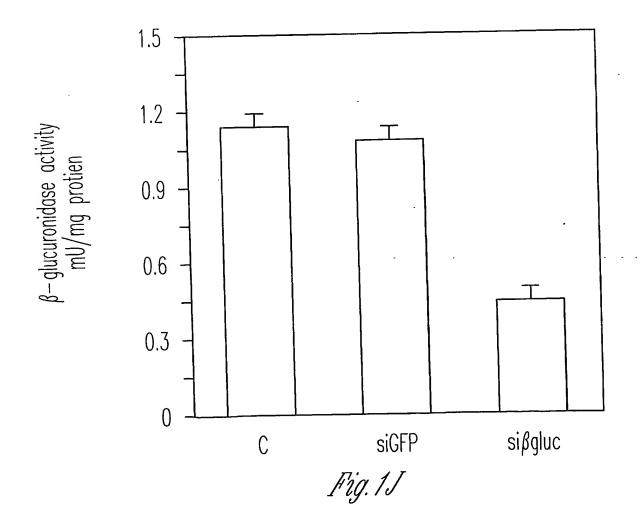


Fig. 11



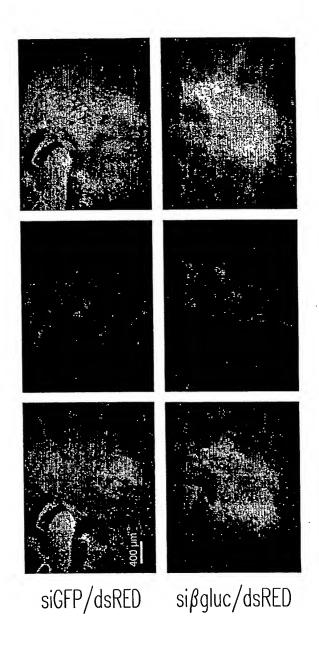


Fig. 24

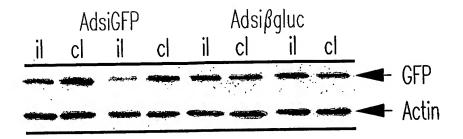


Fig.2B

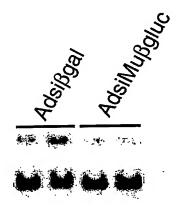


Fig. 20

ig.30

9

siGFP siβgluc

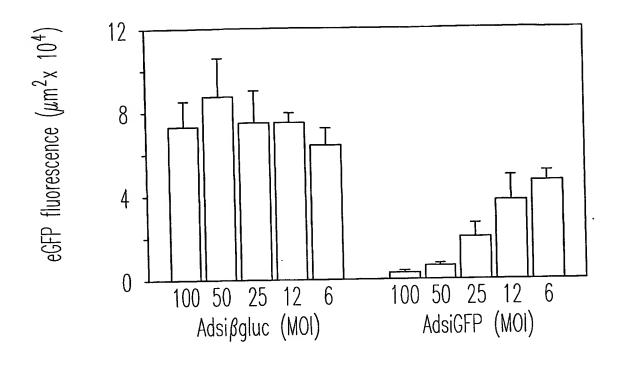


Fig.3D



Fig. 4A



Fig. 4B

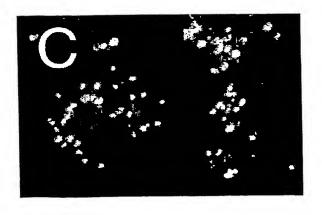
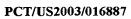


Fig. 4C



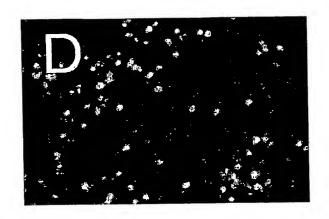


Fig. 4D

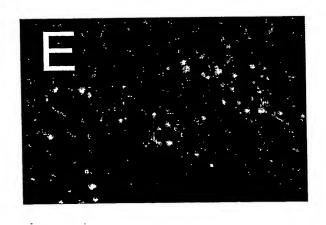


Fig. 4E

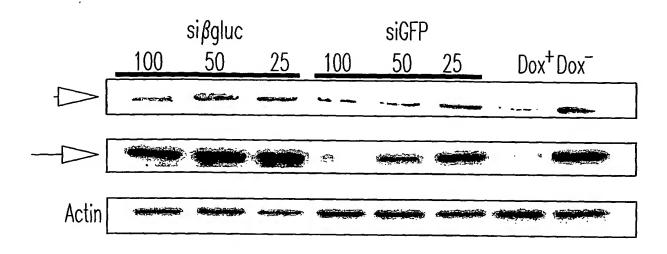


Fig. 4F

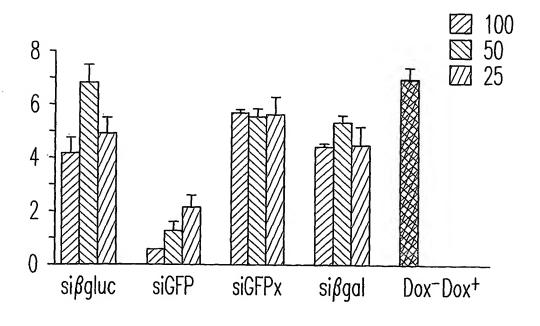


Fig. 4G

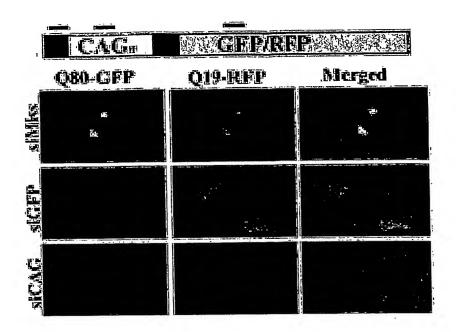


FIG. 5A

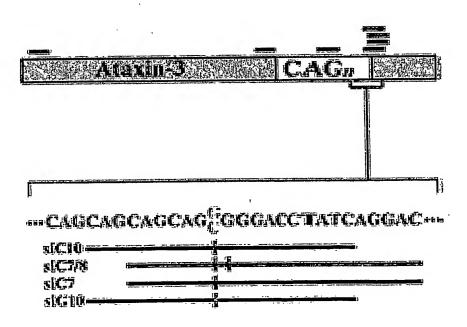


FIG. 5B

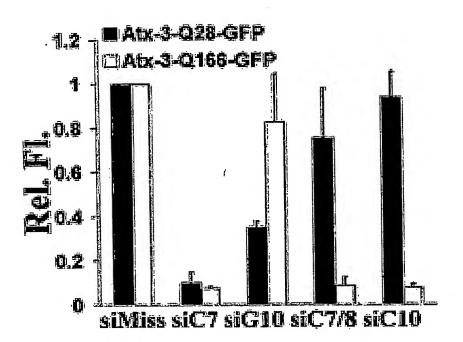


FIG. 5C

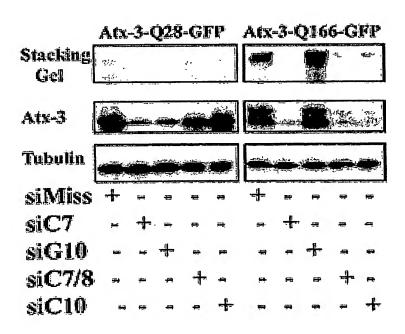


FIG. 5D

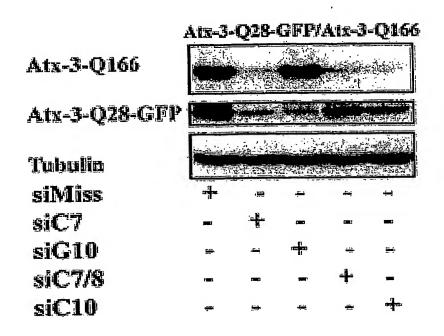


FIG. 5E

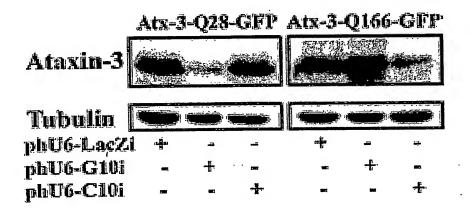


FIG. 5F

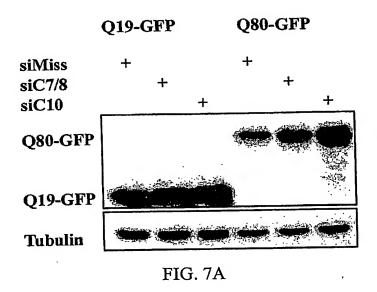
Ataxin-3

Tau

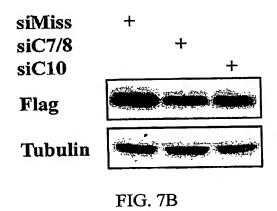
NAME	Primer Sequence (5'-3') NAME	Primer Sequence (5'-3')
siMiss	CGGCAAGCTGCGCATGAAGTTCBEQIDOD:(I) ain '-tau Atgaacttcatgctcagcttgc(5e QIDM);(3)	TCGAAGTGATGGAAGATCACGC(\$\frac{1}{2}\text{b} \nub.; (?\frac{1}{2}) CAGCGTGATCTTCCATCACTTC (\$\frac{1}{2}\text{b} \nub.; (\frac{1}{2}\text{b})
siGFP	ATGAACTTCAGGGTCAGCTTGC(SEQ10Nb:13)si272 CGGCAAGCTGACCCTGAAGTTC(5EQ16Nb:14)	Cagcoggagtogggaaggtgo (Se Ulu PC) Ctgcaccttooggactooggo (Seu Ulu Sa)
siC7	Cagcagcggacctatcaggac (SEQ 1916;15) CTGTCCTGATAGGTCCCGCTGC (SEQ1916;10)	ACGTCCTCGGCGGCGGCAGTGTGC (SEQ)(DN);3) TTGCACACTGCCGCGCGCGGAC (SEQ)(DN);3)
siG10	CAGCAGCAGGGGACCTATC (SEQ\PND;IR) s1406 CTGATAGGTCCCCCTGCTGC (SED\PND;IQ)	ACGTCTCCATGGCATCTCAGC (SE 41 PND: 33) TTGCTGAGATGCCATGGAGAC (SE 610 06:34)
siC7/8	Cagcagccgacctatcaggac(SE01000:19b1a9 Ctgtcctgataggtccggctgc(SE01000:00)	GTGGCCAGATGGAAGTAAAATC(5EQ IVNU:35) CAGATTTACTTCCATCTGGCC (5EQ IVNU:34)
siC10	Cagcagcagcggacctatc (SEQ IDN 6: 2) ia9/c8 Ctgataggtoccgctgctgc(SEQ ID N):2)	GTGGCCACATGGAAGTAAAATC(\$EQIMOI) CAGATTTTACTTCCATGTGGCC(\$EQIMOI)
sin'CAG	TTGAAAACAGCAGCAAAAGC (\$\fa\)\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	GTGGCCAGATGCAAGTAAAATC(SE & 1046;39) CAGATTTTACTTGCATCTGGCC(SE & 1046;40)
siCAG	Cagcagcagcagcagcagc(5& & 1006:25) Ctgctgctgctgctgctgctgc(5&0 1006:25)	

FIG. 6





Flag-Ataxin-1-Q30



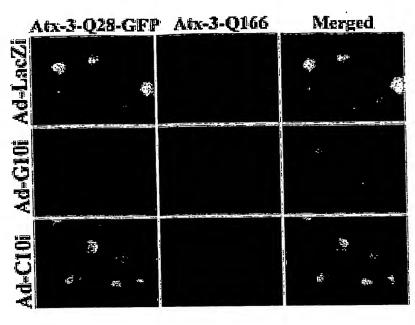


FIG. 8A

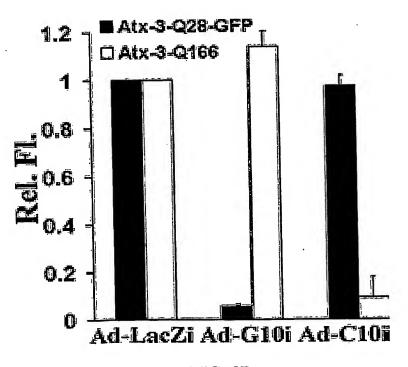


FIG. 8B

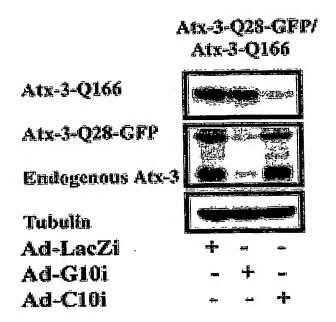


FIG. 8C

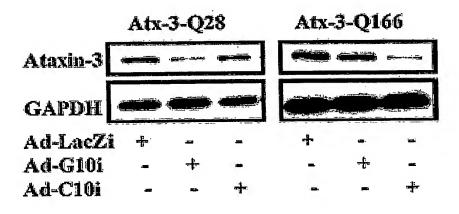
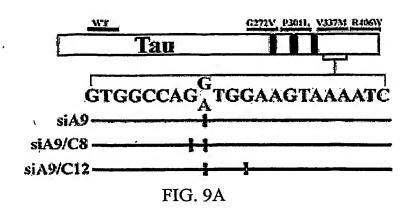
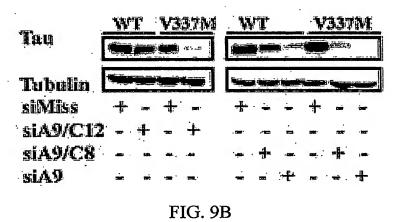
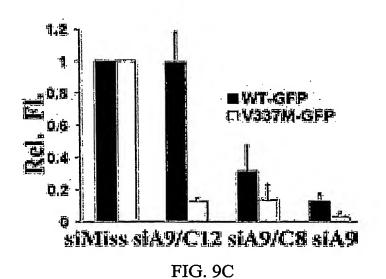
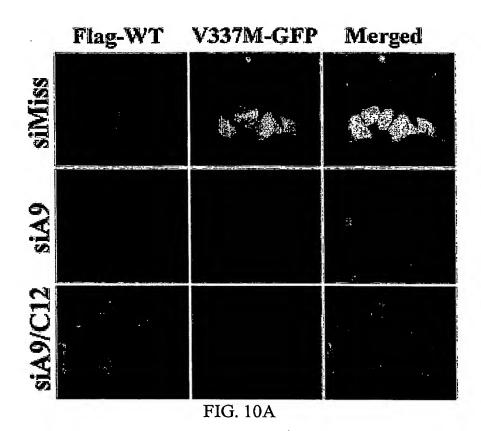


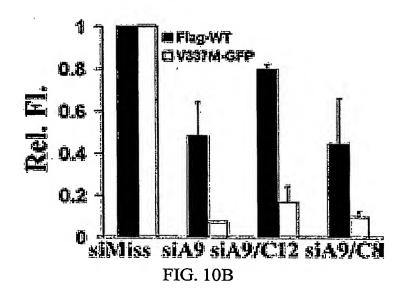
FIG. 8D











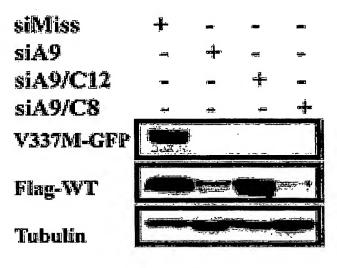


FIG. 10C

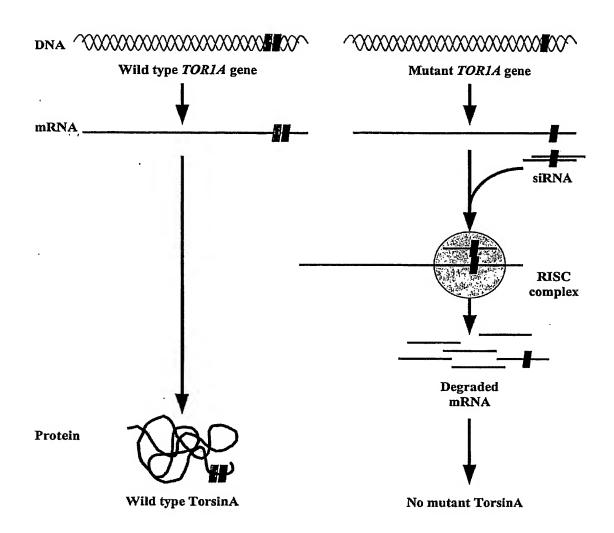


FIG. 11

931 GTAAGCAGAGTGGCTGAGGAGATGACATTTTTCCCCAAAGAG 969 \(\SEG\ \DOO:\4\L) (5EG 1000:44) 3' CTCTACTGTAAAAGGGGTT 5' (SECNO UO:52) GAGTGGCTGAGATGACATTTTTC 3' 960 (SEQ (DNO; 53) 5' CIGAGAIGACAITITICCCC 3' 963 (SECLIDIUS, 51) 3' CACCGACTCTACTGTAAAAGGG 5' CSEG [DNO: 54] 956 (SEG 1000:47) 95365E (210100: 49) 3' CTCACCGACTCCTCTACTGTG 5' (SEQ 10 NO: 48) 3' CTCACCGACTCTACTGTA 5' (SEG ID No. 59) 3'CTTGAAGTACGAGTCGAACGGC 5' (SEO 100043) 5' ATGAACTTCATGCTCAGCTTGC 3' (SE& 10 NO:42) 3' CACCGAAGACCGTGTCGTCGAA 5' 789 5'CAGIGGCIICIGGCACAGCAGC 3' 5'CAGAGTGGCTGAGATGAC 3' 5' CAGAGTGGCTGAGGAGATGAC 3' AWATOTSTINATOTSTA 938 5' 936 mutA-si mutB-si mutC-si com-si mis-si

FIG. 13

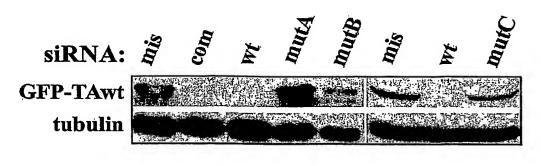


FIG. 13A

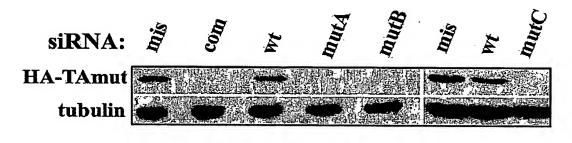


FIG. 13B

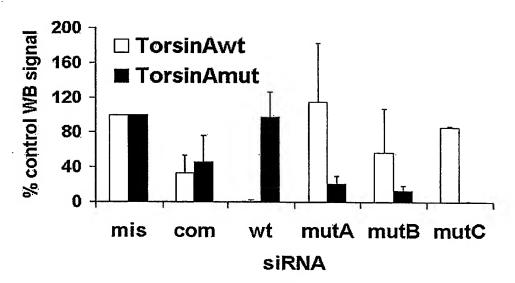
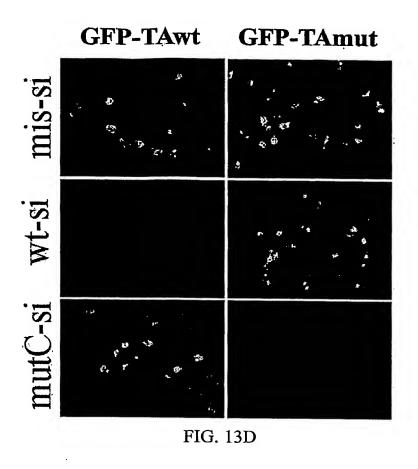
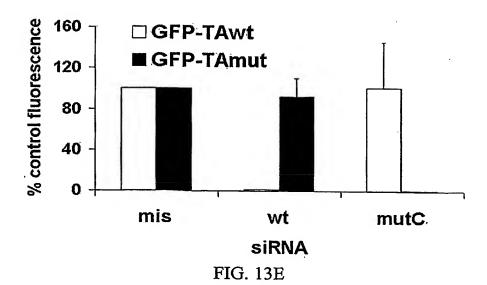


FIG. 13C





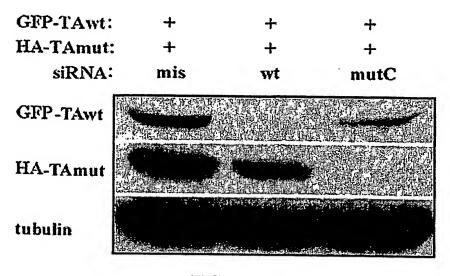
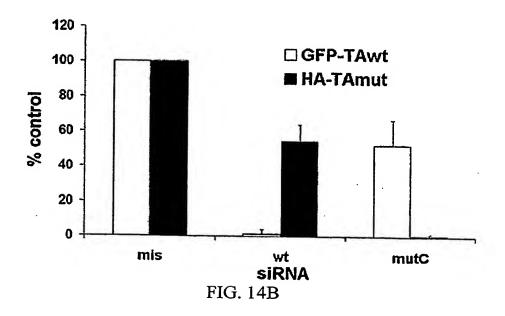


FIG. 14A



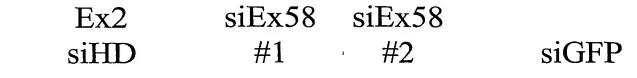




Figure 15

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